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LARGE-SCALE cell preparation of bacterial genomic DNA

1. Grow 100 ml culture of bacterial strain to saturation.
2. Pellet cells for 10 min at 2000 rpm. (use GSA rotor)
3. Resuspend cells gently in 9.5 ml TE buffer. Add 0.5 ml of 10% SDS & 50 μ l of 20 mg/ml proteinase K. Mix thoroughly and incubate 1 hr at 37°C.
4. Add 1.8 ml of 5M NaCl and mix thoroughly.
5. Add 1.5 ml β CTAB/NaCl solution. Mix thoroughly and incubate 20 min at 65°C. (can stay for a moment)
6. Add an equal volume (13 ml) of chloroform/isoamyl alcohol. Extract thoroughly. Spin 6 min at 7000 rpm (use SS34 ~~room temperature~~, to separate phases).
7. Transfer aqueous supernatant to a fresh tube using a wide-bore pipet.
8. Add an phenol / chloroform / isoamyl alcohol. Extraction the ~~equal volume~~
Repeat three times.
9. Add an equal volume of chloroform / isoamyl alcohol.
10. Add 0.6 volume isopropanol and mix. (-20°C overnight)
(-20°C can help to precipitate.)

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Continue:

11. Transfer the precipitate to ~~water~~ 1 ml of 70% ethanol in a fresh tube (just brown-red one), by hooking it on the end of a Pasteur pipet that has been bent (by heating) and sealed.
12. Drying the pellet (just in the air). Remove ~~stuff~~ white and stringy DNA to the ~~1 ml~~ TE buffer.
13. Do the electrophoresis of nuclei and (there are a lot of DNA and RNAs)
14. Use RNase to digest the genomic DNA overnight.

Genomic DNA (continue)

1. Use EcoR I to do restriction, put [genomic DNA 1 ml
 buffer (H) 9 ml
 EcoR I 4 ml] overnight.
- Digest genomic DNA
2. do the electrophoresis at 60 V
 put 10 μ l. digested DNA (90 μ l. digested DNA
 1 μ l dye)
- the result is not good.

Extracting Plasmid DNA of E.coli. (Qiagen Midi and Maxi Pkt.)

- Inoculate a 5-mL LB medium with E. coli. 37°C. shake to afternoon
- Inoculate 200ml medium. Grow at 37°C ~~for~~ overnight

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Extracting plasmid DNA. (continue)

3. Harvest the bacterial cells by centrifugation at $6000g$ for 15min at 4°C .
4. Resuspend the bacterial pellet in 4ml of buffer P1.
5. Add 4ml of buffer p2, mix gently but thoroughly by inverting 4-6 times, and incubate at room temperature for 5 min.
6. Add 4ml of chilled buffer P3, mix immediately but gently by inverting 4-6 times, and incubate on ice for 15 min.
7. Centrifuge at $20000g$ for 30 min at 4°C . Remove supernatant containing plasmid DNA promptly.
8. Re-centrifuge the supernatant at $\geq 20,000 \times g$ for 15 min at 4°C . Remove supernatant containing plasmid DNA promptly. Alternatively,
~~the sample can be fill~~
9. Equilibrate a QIAGEN-tip 100 by applying 4ml Buffer QBT, and allow the column to empty by gravity flow.
10. Apply the supernatant from step 8 to the QIAGEN-tip and allow it to enter the resin by gravity flow.
11. Wash the QIAGEN-tip with 10ml (two times) buffer QC.
12. Elute DNA with 5ml buffer QF.
13. precipitate DNA by adding 3.5ml room-temperature isopropanol to the eluted DNA. Mix and Centrifuge immediately at $13,000 \times g$ for 3.5min at 4°C . Carefully decant the supernatant.
14. Wash DNA pellet with 2ml of room-temperature 70% ethanol, and centrifuge at $13,000 \times g$ for 10min. Carefully decant the supernatant without

disturbing the ~~pette~~ pellet.

15 Air-dry the pellet for 5-10 min. and redissolve the DNA in a suitable volume of buffer \rightarrow (TE buffer \rightarrow 1/2 ml)

16 Do electrophoresis to check it. \rightarrow to digest? \rightarrow next pg

8/3/99

High-efficiency transformation by electroporation

1. Inoculate a single colony of E. coli cells into 5 ml LB medium. Grow shaker overnight at 37°C with moderate shaking.

2. Inoculate 2 ml of the culture into 50 ml LB medium in a sterilized flask. Grow at 37°C shaking at ~ 1000 rpm. (if > 2 hr more time)

3. Chill cells in an ice-water bath 10 to 15 min and transfer to a prechilled 50ml centrifuge bottle.

4. Centrifuge cells 10 min at 4200 rpm. 2°C

5. Pour off supernatant and resuspend the pellet in ice-cold water. Add 25ml ice-cold water and mix well at 10 min 8000 rpm. repeat 2 times. Centrifuge cells as in step 4. (throw away the supernatant to the bleach) (bottle including)

6. Pour off supernatant immediately and resuspend the pellet by swirling in remaining liquid.

7. Add another ml ice-cold water, mix well, and centrifuge again in step 4

8. Pour off supernatant immediately and resuspend the pellet by swirling in remaining liquid.

9. Put remaining liquid to the eppendorf tube. centrifuge it at 12000 rpm 5 min. pour off by using vacuum

10. add 100ul water to the tube. mix well. put it on ice.

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- 10 Set the electroporation apparatus to 2.5 kV, 25 μF. Set the pulse controller to 200 or 400 ohms? (Under the machine instruction) and 20μL bacterial
- 11 Add 1μL plasmid DNA (PGP 704) in to tubes containing fresh or thawed cells (on ice). Mix by tapping the tube or by swirling the cells with the pipettor.
- 12 Transfer the DNA and cells into a cuvette that has been chilled 5 min on ice. Shake slightly to settle the cells to the bottom, and wipe the ice and water from the cuvette with a Kimwipe between
- 13 Place the cuvette into the sample chamber.
- 14 Apply the pulse by pushing the button of flipping the switch.
- 15 Remove the cuvette. Immediately add 50 μL media and transfer to a stir culture tube with a Pasteur pipettor. Incubate ~~at 37°C~~ 60 min with moderate shaking including LB broth.
- 16 Plate aliquots of the transformation culture on LB plates containing antibiotics.

plasmid digest

100 μL 08/03/99

| | | |
|---------------|--------|------------------------------|
| PGP 704 | 1 μL | |
| 10x buffer | 1 μL | |
| DNasefree H2O | 7.5 μL | |
| Zerk I | 0.5 μL | 37°C 2 hr. → 65°C for 15 min |
| | | run 0.1 mL on gel |

dilution 10⁻⁶ 100 μL to LB DAP10⁻¹ 10⁻² 10⁻³ 10⁻⁴ 10⁻⁵ 10⁻⁶ round to LB DAP DAP.

result

10⁻¹ 10⁻² zero 236 v10⁻³ 100 colonies

DB electrophoresis.

1. 15 μl marker (1kb)
2. 65 μl digested DNA + 13 μl 6x dye. first line
Second line.
3. Cut out 7-10 kb gene.
4. elute into Bio-rad ~ promega.

Fragment Isolation Protocol from NuSieve Gel.

1. Place the electrophoresed gel on the UV transilluminator and view under long wave light. Locate the fragment of 7-10 kb and cut out the band using a glass cover slip and place the gel fragment into a 1.5ml microcentrifuge tube.
 2. Heat to 65°C until gel slice melts
 3. Add 3 volumes of BioRad DNA isolation binding buffer and for every 2 μg of DNA add 5 μl of matrix
 4. Attach a clean 3cc syringe to a promega spin column and add the solution to it.
 5. Push this slowly through the column (1 drop/3 sec)
 6. Detach column from syringe and remove plunger.
 7. Wash with 2 mls of 95% isopropanol.
 8. Spin at 12,000 rpm for 20 sec.
 9. Dry Column in Hybridization oven for a few minutes at 68°C
 10. Add 30-50 μl of water or TE. wait 1 min. and spin again at 12,000 rpm for 20 sec.
 11. Run 10 μl of the eluant on a gel to check recovery.
- 5 μl

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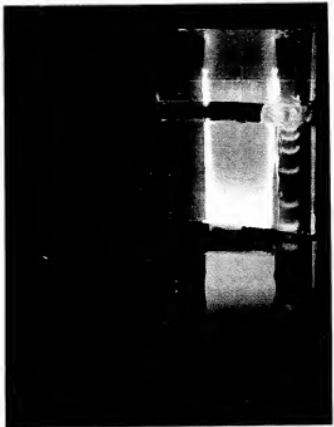
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4 8

Notebook Number:

Date: 08/04/99



dig RFLP 2.69 contac (mb)

cut out 7-10 kb

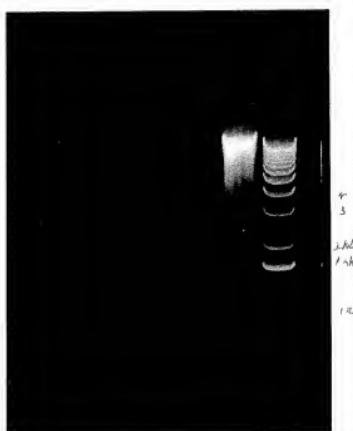
Do electrophoresis to check the result.

DNA. 5ul. + 1ul dye.

15ul Marker

→ 60v. min

Results:



Conjugation:

1. Donor Strain (E. coli MGN 617 PGP704 or MGN 617 PLOF Km) are grown overnight with shaking at 37°C in 2 ml LB broth containing 200 µg/ml ampicillin and 50 µg/ml Kanamycin. (DAP)
2. Grown P. multocida 11039 overnight with shaking at 37°C.
3. 50 µl MGN 617 or 50 µl MGN 617 + 100 µl 11039 to 5 ml of 10mM MgSO₄.
4. Vortex for a few seconds.
5. Transfer to a 5ml disposable syringe and filter through a 25 mm Milipore filter
6. Drain the filter and carefully remove it from the filter case with sterile forceps (curved tip work best)

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10 $\text{LbApDAP} \rightarrow \text{LbAp}$

100ul TPG

200ul

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+ 50ul DAP? Date: 08/05/99

7. Place the filter on a LB plate cell side up.

(avoid bubbles between the filter and the agar)

PLOF

8. Incubate at 37°C for 8-18 hrs.

2.36V - Km

08/06/99

Results 9. Resuspend the filter using 3ml 10mM MgSO₄. Vortex.

| | | | |
|---------|-------------|------------------------|------------|
| PGP 704 | BH1 Ap | 100ul 10 ⁻¹ | } negative |
| | BH2 ApK | 100ul 10 ⁻¹ | |
| | BH2 | 100ul 10 ⁻⁵ | |
| PLOF Km | LB ApDAP | 100ul 10 ⁻⁵ | → negative |
| | BH1 ApKm | 100ul 10 ⁻¹ | |
| | BH2 Km | 100ul 10 ⁻¹ | |
| | BH2 | 100ul 10 ⁻⁵ | |
| | LB ApKm DAP | 100ul 10 ⁻⁵ | |
| | | 200ul | |

Transformation by electroporation

PGP 704:

PLOF Km 10⁻¹ 10⁻¹ 10⁻² deletion

Results



08/09/99

Digestion:

93-146 genomic DNA (cc plasmid prep.)

Leave incubate at 37°C 30min

Restriction Digestion

Z-EcoRI

DNA 51ul

10x buffer 6

Z-EcoRI 3

37°C 4h → put in 65°C water bath

08/10/99

3b electrophoresis

903-146 genomic DNA (after digest) 60 uL
dye (5X) 10 uL

↓
run gel (marker 1kb 12 uL)

Fragment Isolation Protocol from Gel (See Page 7)

1. Cut out 7-10 kb and place the gel fragment into microcentrifuge tube.
2. Heat to 65°C until gel slice melts.
3. Add 3 volumes (630 uL) of BioRad DNA isolation binding buffer ~~exclusion~~ (1 mg \approx 1 uL volume) and usually add 30^{uL} matrix.
4. Attach a clean 3cc syringe to a Promega spin column and add the solution to it.
5. Push this slowly through the column.
6. Detach column from syringe and remove plunger.
7. Wash with 2 mls of 95% isopropanol.
8. Spin at 12,000 rpm for 20 sec.
9. Dry Column in Hybridization oven for a few minutes.
10. Add 30 uL of TE wait 1 min and spin again at 12,000 rpm for 20 sec.
11. Run Gel.

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9130199

Digest:

P1039 6ml
Buffer 7ml
EcoRI 3ml

37°C 2 h / 65°C 15 min.

9130199

1. Run a gel. 7ml.
2. Cut out 7-10kb.
3. Use another kit to extract DNA.

QIA quick Gel Extraction kit protocol (another kit)

1. Excise the DNA fragment from the agarose gel with a clean, sharp scalpel.
2. Weight the gel slice in a tube. Add 3 volumes of Buffer QG to 1 volume of gel (100mg ~100ul).
3. Incubate at 50°C for 10min.
4. After the gel slice has dissolved completely, check that the color of mixture is yellow.
5. Add 1 gel volume of isopropanol to the sample and mix.
6. To bind DNA, pipet the sample onto the QIA quick column and apply vacuum. After the sample has passed through the column, switch off vacuum source.

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Run gel (Southern Blotting)

1. Digest E. coli Sal I Bam HI Xba I 11039E. coli \Rightarrow in back

2. Rinse & Rinses

3. photograph with a ruler laid alongside the gel so that band positions can later be identified on the membrane.
3. Rinse the gel in distilled water and place in a clean glass dish containing ~ 10 gel volumes of c. 25 M HCl. Shake slowly on a platform shaker for 30 min at room temperature.
4. Pour off the HCl and rinse the gel with distilled water. Add ~10 vol denaturation solution and shake as before for 30 min to 1 hr.
5. Pour off the denaturation solution and rinse the gel with distilled water. Add ~10 vol neutralization solution. Shake as before for 30 min to 1 hr.

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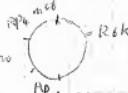
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11/19/99

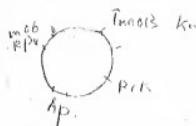
Conjugation:

See Pg 10;

put the conjugation in

(1) MGN 617 10^{-1} BHI Km(2) 10^{-1} BHI Ap-(3) MGN 617 PGP 10^{-1} BHI Ap Km-
704 10^{-1} BHI Km 10^{-1} BHI Ap → ran gro 10^{-1} BHI Ap Km

(4) MGN 617 PLOF Km

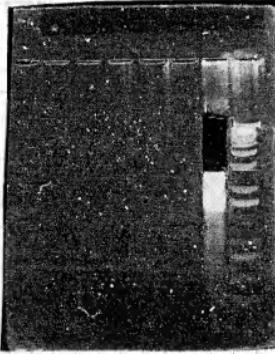
 10^{-1} BHI Km K2 (100 uL. 20-50 colonies) 10^{-1} BHI Ap 10^{-1} BHI Ap Km100 uL 10^{-5} BHI 11039 10^{-5} LB DAP/pkm → MGN 617↓
200 uL DAP

11/23/99

Cut out 2-6 Kb,
7-10 Kb, and more than
10 Kb

use Qiaquick gel
extraction kit to extract
DNA.

result:



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the digest is not

In P_EEM-38 has not been
digested completely.

So there are three bands.
(supercoiled DNA and one
strand DNA)

So repeat the digest

10039 having 1 week long can not grow in BH2 broth
but can grow in BH2 plate

Inoculate 10039 and 1004-617. PGM 704. PLOTTING again.

Do digest again 2.0 μl DNA (P_EEM-38)
4 μl ZEOL I + 3 ml buffer
over 2 h. + 15° 65° C

11/13/2000

Sep 1 μl + 4 μl buffer → 37° C 2 h. → 15° 65° C
+ ~~15°~~ 20° C 10 min ECHO + 4 μl 3M NaAc → fraser
+ ~~70°~~ 10 min 13500 rpm 15' → pour off supernatant →
100 μl 2.70% ECHO 13500 rpm 10' → pour off supernatant
→ dry → run a gel (1 μl)

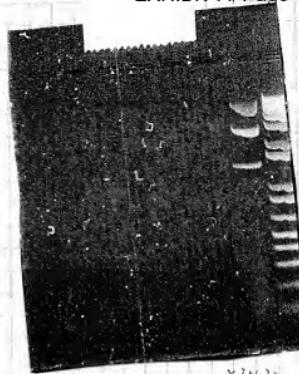
Conjugation LB plate + 200 μl DIP + 100 μl RPTG.
do conjugation

BH2 → BH2 + 100 μl Ap → BH2Ap
BH2 + 100 μl Km → BH2Km
BH2 + 100 μl Ap + 100 μl Km → BH2ApKm

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2/4/00

Run a gel Digest. 21.5 mL DNA
2.5 mL buffer overnight (37°C)
1 mL EtOH
→ 65°C 15'

Run a gel. Cut out 2-8 kb, 6-10 kb monomers
(because the extraction DNA ^{was} always bigger than
it is. So cut out ^{was} _{little} smaller band)

2/7/00

Extract DNA. (use Qiaquick Spin Kit)

2/8/00

Run a gel. but no DNA

improve: Clean DNA before

kit

2/10/00

* Clean DNA

1/2 volume 3M Na acetate

2 vol 100% eth

-20°C > 1 h.

centrifuge > 13,000 rpm 4°C 15'

Add 10.0 mL 100% EtOH
2 mL
Re-spin
Dry

Remove supernatant

Add 100 uL 100% EtOH

centrifuge > 13,000 rpm 4°C
10 min

Remove supernatant dry ^{air} ~ 5 min

Resuspend 30 uL DNase free H2O

Run a gel → cut out

2/13/00

- Run a gel (genomic DNA 11039)
 Do digest

It just has a little DNA
 may because the DNA doesn't
 seem dissolve well.

2/14/00

Set up PCR reaction

| | ① | ② | ③ | ④ |
|------------------|------|------|------|------|
| primer SC 1011 | 0.3 | | 0.3 | 0.3 |
| primer SC 1012 | 0.8 | | 0.8 | 0.8 |
| dNTP | 4 | 4 | 0 | 4 |
| 3.3X buffer | 6 | 6 | 6 | 6 |
| Mg | 2.4 | 2.4 | 2.4 | 2.4 |
| H ₂ O | 6 | 7.6 | 10 | 6 |
| | 20ul | 20ul | 20ul | 20ul |

higher layer

| | ① | ② | ③ | ④ |
|------------------|------|----|----|------|
| control DNA | 0 | 0 | 0 | 0.5 |
| 3.3X buffer | 9 | 9 | 9 | 9 |
| enzyme | 1 | 1 | 1 | 1 |
| H ₂ O | 20 | 20 | 20 | 19.5 |
| | 30ul | | | |

put 11039 in 56° for 2 hrs because a lot of
 DNA do not dissolve in TE buffer.
 more than

Run a gel (11039 genomic DNA .

① Line 1 μl DNA)
 ② Line 3 μl DNA .

Do digest (11039) 7.5 μl total

| | |
|---------------|-----------------|
| 62.5 μl DNA | } 3/1 overnight |
| 7.5 μl buffer | |
| 5 μl Enzyme | |

65° 15'



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in 3 sq rubber mat dishes put 2 pes chromatograph
filter paper (this is too soak up buffers.)

- 1) Denaturation solution - takes very little just enough
to soak up filter. 3 min.
- 2) Neutralization buffer 5 min
- 3) 2x SSC 5 min. place it on the paper and dry for 30'
cross link on program c3 → put them in plastic.

4113100.

1. preheat the hybridization buffer to 42°C
2. In a suitable container prewet the blot in 5xSSC. Loosely
roll the blot and place inside the tube. Add small amount
of 5xSSC to the tube and "unroll" the blot ensuring no
air bubbles are trapped between the membrane and the
tube. Don't allow the blot to overlap itself
3. Pour off the 5xSSC and add the appropriate volume
of ~~the~~ hybridization (20ml). buffer.
4. Prehybridize in oven for at least 30 min at 42°C
5. Prepare the labelled nucleic acid probe as instruction
P13.
6. Dilute the DNA to be labelled to a concentration of 10 ng/ml
using the water supplied.
7. Denature 10ng of the DNA sample (10ul) by heating for
5 min in a boiling water bath. (13ul dian + 7ul H₂O)
8. Immediately cool the DNA on ice for 1 min. Spin briefly
in a microcentrifuge to collect the contents at the
bottom of the tube.
9. Add an equivalent volume of DNA labelling reagent (10ul)
to the cooled DNA. Mix gently but thoroughly

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5/1/00

Prepare sequencing for genomic DNA.

Set up sequence program. (Lang sequence)

Big dye: 2⁻3mer primer purified gen DNA H2O
 Roe-Lab (Pmdes) 1039 base sul
 rx reaction 8ul (ul 13pmol ul)

total 20ul

p.s. primer (add 3ul primer to 97ul H2O)

HADAM P2

HADAM M2

PCR

5/1/00 2 prepare sequencing

1. Column Hydration

- ① Remove the top of column, then add 0.8ml of reagent grade water. invert for a few times. make sure no bubbles. leave the column for at least 2 hrs. at room temp.

② Removal of Interstitial Fluid:

- ① remove the top cap first. then remove the end stopper from bottom
- ② allow excess column fluid to drain into a wash tube. discard this fluid.
- ③ spin the column and wash tube in centrifuge at 750g for 2 mins. discard it

3. Sample processing:

- ④ Add 20ul sequenced PCR product into tube column. make sure it's be into the column, but don't touch the column.
- ⑤ spin the column at 750g for 2 min - the purified sample will collect in the collection tube.
- ⑥ Dry the sample in a vacuum centrifuge.

4. ① vortex resuspend sample in 25ul of template reagent in

- ② vortex 30sec } X2 2min denature in 95°C quick spin }
- Chill 5 min

put read in sequencing tube
 freeze & resuming later if needed
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Signed . . . sequencing didn't work . . . Signed

11800 foci/l $\Rightarrow 1780000 \text{ colonies / ml}$

$$\frac{1.78 \times 10^5}{0.005 \text{ ug}} = 3.56 \times 10^7 \text{ / mg DNA}$$

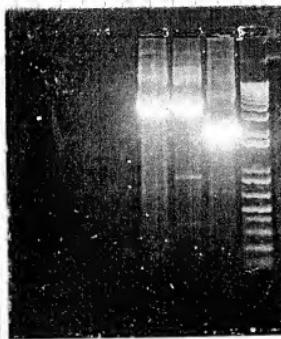
Result: the problem is Xgal. the old Xgal is not useful for producing blue colonies
do electroporation again, and inoculate the restricted bacteria to the LB Broth

6/13/00

Extract the plasmid DNA (restricted Baet)
put the electroporation to refrigerator

Run a gel (plasmid DNA Φ \emptyset \odot \ominus)

\downarrow
2ul + 2ul + 1ul dye
DNA H2O



\odot \ominus \emptyset

\odot and \ominus is definitely inserted. Do the digest.

20ul. total

1ul DNA

2ul buffer

0.5 Elect

16.5ul H2O

overnight 37°C load 3-4ul on a gel

6/14/00

do electroporation, southern hybridization:

1. Run a gel (plasmid DNA Elect digested (20ul))

load 3-4 ul per well.

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★ 2x SSC 5 min x2 at room temp.
 5x SSC 0.5% SDS 42°C 30' 热, oven
~~15°C~~ 50 ml ~~50°C~~ 2nd 10°C
 2x SSC 用软纸 (Kimwipes) 将膜吸干并擦.
 >x SSC 洗干净 rinse and shake 5' ~~14°C~~

7/9/00

Do ligation.

1 ml PGIEM-32 EGRI + Sip tit.
 8 ml insertor (1039 3-5 kb extraction)
 1 ml buffer
 0.5 ml ligase (New England Lab)
16°C overnight

7/10/00

65°C 15'

Do electroporation.

Note: cuvette should be chilled 5 min on ice
 immediately add 1 ml LB medium and transfer
 to a sterile culture tube.

12:100

Run a longer gel for 11039 ECR I.
 1 lane: ladder (4ml) 5830 pm. 20v.
 3 lanes: 11039 ECR I. 5 - ladder (4ml)

It goes very slowly. I should run this gel at 30v.

12:45 end.
 1:10 continue
 6:10 em.

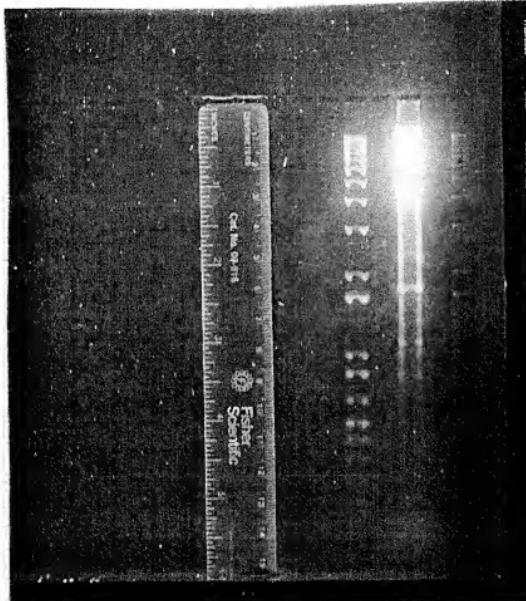
gel size 6x15

paper towel 8x15

filter paper 3 (^{5.5}_{5.5} x 14)

filter paper 5 (8x15)

membrane 7x14



the part from 6kb to 8kb has been reseparated Dr
 Southern hybridization.

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make hybridization buffer.

water buffer

5 g blotting reagent
overnight at 27°C.

room temp. stir for 1 hr.

then 42°C at least 30'.

8/10/100

Continue Southern hybridization and Sequencing.

8/10/102

Sequencing didn't work again, but Southern hybridization did work.

set up the PCR reaction to make more 1-10-2 and check the plasmid DNA.

primer dilution

95 uL H₂O

2.6 uL Taq polymerase

2.6 uL - - - P1

100.0 uL

→ 1 mL

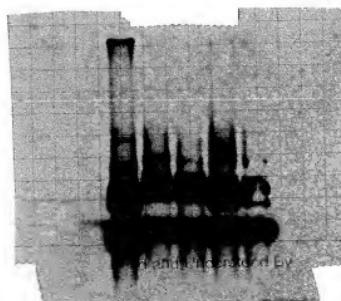
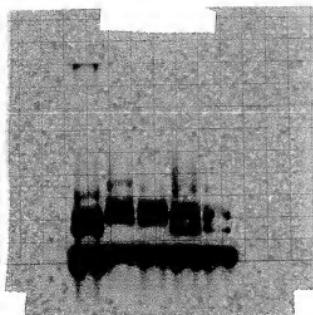
PCR reaction

| | | |
|---|------------------|------|
| ① | 1-10-2 | 0.5 |
| | dNTP | 5 |
| | primers | 1 |
| | buffer | 5 |
| | Taq | 0.5 |
| | H ₂ O | 38.5 |

| | | |
|---|------------------|------|
| ② | plasmid | 1 mL |
| | dNTP | 5 |
| | primer | 1 |
| | buffer | 5 |
| | Taq | 0.5 |
| | H ₂ O | 37.5 |

| | | |
|---|------------------|------|
| ③ | 1-8-1 | 2 |
| | dNTP | 5 |
| | primers | 1 |
| | buffer | 5 |
| | Taq | 0.5 |
| | H ₂ O | 36.5 |

| | | |
|---|------------------|------|
| ④ | Control | 0 |
| | dNTP | 5 |
| | primer | 1 |
| | buffer | 5 |
| | Taq | 0.5 |
| | H ₂ O | 38.5 |



Date: _____

8/25/00

Run a sequencing reaction

| | |
|------------------|------|
| bigr dye | 1 ml |
| plasmid | 2 ml |
| primer | 2 ml |
| H ₂ O | 8 ml |

Run a gel for genomic DNA
 (0.5%) in order to separate
 of ladder

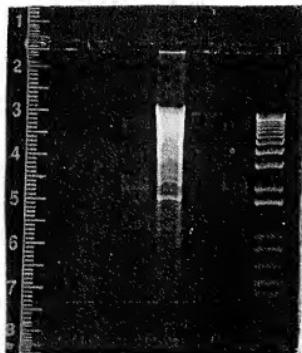
Should keep
 some a little
 11039 Hind III
 Left for Southern in
 V-region

8/28/00

Run a gel. 1 ml for extracted DNA
 1 ml for ECR I 11039 digest & positive control)

- ① extracted min 4-6 kb
- ② " 6-8 kb
- ③ 11039 ECR I
- ④ repeat ② → another tube
- ⑤ repeat ④ → another tube

8/29/00



the size of both tubes is bigger
 than needed. So extract DNA
 from 3-4 gel slice.

8/30/00

Run a gel

10/9/00

Digest - clone -> a plasmid DNA to get probe

| | |
|--------------------|-------------|
| 3 ul | 3 ul DNA |
| Digest | 3 ul buffer |
| <u>1 ul EcoR I</u> | |
| <u>2 ml H2O</u> | |

30 ul → 2 h. 37°C

15' 65°C

cut out from 1.4 kb.

10/10/00

PGEM-3Z HindIII.

1. Continue plasmid DNA + sap treatment
2. do gel extraction.

10/11/00

Run a gel for gel extraction (for probe) and PGEM-3Z HindIII + sap trt. (both are same) gel extraction (probe), just need to use same for detect.



10/12/00

the electroporation didn't work. try electroporation. 1 ul PGEM3Z +

10^{-2} . 10^{-8} .

it did work.

Read and Understood By

Signed

Signed

1/31/01

Do PCR again

EXHIBIT A, Page 27

| | digest | (1) | (2) | (3) |
|-------------------|--------------|------|------|-----|
| | DNA | 1 | 1 | 1 |
| Pr multicoder A:3 | (1) | (2) | (3) | |
| | 0 | 1 | 1 | |
| dNTP | 5 | 5 | 5 | |
| primer | 1 | 1 | 0.5 | |
| buffer | 5 | 5 | 5 | |
| New tag | 0.5 | 0.5 | 0.5 | |
| H ₂ O | 38.5 | 37.5 | 38.0 | |
| | <u>50 ul</u> | | | |

95°C 2 min

95°C 30 sec.

72°C 1 min

4°C hold

1 cycle

25 cycles

2/10/01

Do PCR again. Add more genomic DNA and increase annealing temperature to 66°C.

there's
a little DNA on
the gel. It's
not clear. So
Add more genomic
DNA to do PCR
again



| | (1) | (2) | (3) | (4) |
|------------------|--------------|------|------|------|
| K mult (A:3) | 0 | 3 | 6 | 10 |
| dNTP | 5 | 5 | 5 | 5 |
| primer | 1 | 1 | 1 | 1 |
| buffer | 5 | 5 | 5 | 5 |
| Taq | 0.5 | 0.5 | 0.5 | 0.5 |
| H ₂ O | 38.5 | 35.5 | 32.5 | 28.5 |
| | <u>50 ul</u> | | | |

annealing

temperature 66°C 1'

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315101

Run a gel for H digest.

Do a Klenow for clone n ~~E~~ claz and T-blunt digest

Klenow : 20ul digest (the concentration of my DNA is high enough)

0.8 uL dNTPs (200ul of each in working stock).

and 0.1 mg/ml BSA (I skip this step)

0.5 uL Klenow

I didn't add buffer because the Multi Core (buffer) is compatible for Klenow

incubate @ 37°C for 30 min. HI @ 75°C for 10 min.

(I lost this Klenow product)

316101

Do digest for B. C. E. F. G.I.H. and run a gel for it.

I select B. C. G to run a seg. react again.

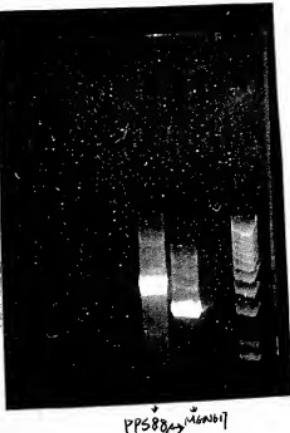
And to make sure which one is right.

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Signed

Signed

BIO TOPIC: Bacteriophage Lambda Phage Lambda DNA Lysogenic Cycle, pL229, pL229, pL229



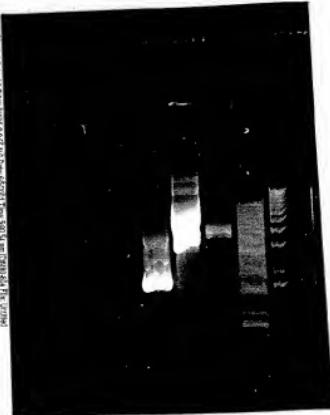
615101.
Do the digest for PPS88 (Xba I + Kpn I)

5 μl DNA
12 μl H₂O
(multicore) buffer 2 μl
0.5 μl XbaI
0.5 μl KpnI
2 h 37°C

inactivate 65°C 15'

Run a gel for
digestion, genomic DNA
PCLPM2.
(from clone A)
and pbluescript plasmid

all the samples → 1 μl



PPS clone M6N.
88 A
plasmid

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do sequencing reaction for
number 5

Ran a gel for Ø Ø Ø

big eye 8 mm EXHIBIT A, Page 30
plasmid 2 ml
primer T3 2 ml
water 8 ml

| Sample | Type | 230nm | 260nm | 280nm | 320nm | ratio | Concentration | Dilution Factor | Purity |
|---------------------------------------|-------|-------|-------|-------|-------|---------|---------------|-----------------|-----------|
| Bluescript and dam ⁺ dsDNA | dsDNA | 3.631 | 5.554 | 3.175 | 0 | 1.74929 | 0.2777 ug/ul | 1 | 50 174.93 |

↓
number 5.

6/29/01

| Sample | Type | 230nm | 260nm | 280nm | 320nm | ratio | Concentration | Dilution Factor | Purity |
|--------|-------|-------|-------|-------|-------|---------|---------------|-----------------|-----------|
| pLS88 | dsDNA | 0.577 | 1.176 | 0.606 | 0 | 1.94059 | 0.0588 ug/ul | 1 | 50 194.06 |

8 ml pLS 88

XbaI / BamH I 0.5 ml

buffer (multicore) 1 ml

water 0.5 ml

Xba I 0.5 ml

BamH I 0.5 ml

multicore 1 ml

7/3/01

PCR:

Template pcyPm3

0.5 ml

1 ml

control

5 ml

5 ml

5 ml

1 ml

1 ml

1 ml

5 ml

5 ml

5 ml

0.5 ml

0.5 ml

0.5 ml

38 ml

37 ml

38.5 ml

508 R102-118 Jun 12 2001

GLNU6YS

T7PR

GLDmM10

G-CATGAGTGTGGCAAGTC

CAGCTTA

5739/ul

5739/ul

48.490

48.490

33.759/ul

33.759/ul

MW=1991

MW=1991

29 nmol

29 nmol

10.300

10.300

11.719/ul

11.719/ul

MW=1972

MW=1972

29 nmol

29 nmol

11.719/ul

8/15/01 Cont.

EXHIBIT A, Page 31

PCR 4 XLB electrooperation...

Blotted plates with hybrid membrane
followed protocol for colony Hybridization

Cross linked Membrane - stored at -20°C ON

8/16/01

Colony Hybridization PCR 4 XLB

made probe using Biol PCR-4 + T4L Heo in ECL Kit
preheat hybridization buffer to 42°C

wet Screen mesh + Membranes
Biol + put in hybridization tube
prehybridize with SSC for 1 hour
 $5X$

use 30mls hybridization buffer

probe Biol 5"

Cool on ice 5"

10μl labelling reagent

10μl glutaraldehyde

10° 37°C

add 1ml hybridization buffer from tube - add buffer +
probe to tube 42°C ON

day 8
8/17/01

Colony Hybridization Cont.

preheat 1° wash buffer to 42°C

discard hybridization buffer. ~~Add 5mls~~

Add 50mls SX SSC return to Oven 5 min

discard SX SSC add 1/3 vol of tube w/ 1° Wash Buff

Return to Oven for 20 min

Wash again in 1° wash Buff for 10 min 2X

remove blots from tube place in glass container

Cover w/2x SSC

Shake @ RT for 5 min X 2

Detection

8ml/membrane 13mls #1 13mls #2 3 membranes
incubate 1 min
place on Saran

Read and Understood By

Signed

Signed

9/19/01 Cont.

Culture reached 260 KU after 4 hrs incubation at
37°C

- (2) Put cultures into 5534 tube
- (3) Incubate on ice 15"
- (4) CF 4°C 5000 xg (6500rpm) 15"
- (5) Wash 2X w/ 1mM HEPES buffer (pH 7.0)
- (6) resuspend pellet in 10% glycerol to 1/2 their
Original Vol = 2 mls
- (7) CF 5000xg 10"
- (8) resuspend in remaining glycerol solution after
Supernatant is decanted
- (9) Flash freeze in EtOH + dry ice bath
- (10) Freeze at -80°C

P mult 11039 is highly encapsulated + forms a very soft pellet
I lost the pellet in the 1st wash step, so more cells will
have to be prepared

Electroporate pLS88 into P.mult 1069

1 μl pLS88 plasmid prep (in plasmid box spot G9)

40μl 1069 comp cells

2.5 KV
200 Ω
25 μF
Tcc 5.12 msec

Incubate 37°C. 1 hr in LB

plate 10⁰, 10⁻¹, 10⁺¹ 100μl on LB strep 37°C

10⁺¹ = after plating 10⁰ + diluting 10⁻¹, et culture + resuspend pellet in
100 μl media - plate 100 μl

RESULTS
10⁺¹ 134 colonies
10⁰ 2
10⁻¹ 0

9/25/01 Tuesday

Pass colony from R6E12 #4 blot - do a plasmid prep
Electroporate pLS88 into P. mult 11039

1 ul pLS88 plasmid prep (in plasmid box spot 59)

40 μl 11039 comp cells (flash frozen in liquid N₂)

2.5 KV

200 Ω

25 μF

Tc = 5.16 msec

Incubate 37°C 1 hr LB

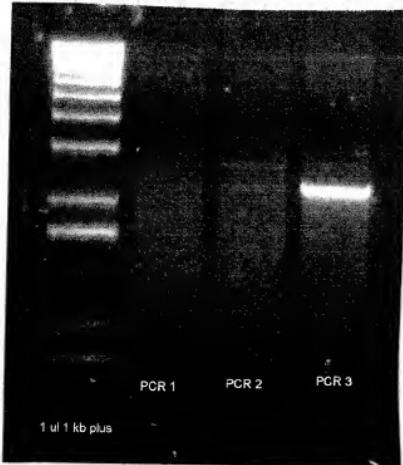
Plate 10⁰, 10⁻¹, 10⁺¹ 100μl on LBstrep 37°C

RESULTS

10⁺¹ 80

10⁰ 13

10⁻¹ 1



See pg 35 for PCR Rx setup

Annealing temp = 48°C

Extension time = 30 sec

PCR Rx did not work again

0.7% agarose 3.5 μl of star
5 μl of PCR Rx loaded

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Signed

Signed

Notebook Number:

Date:

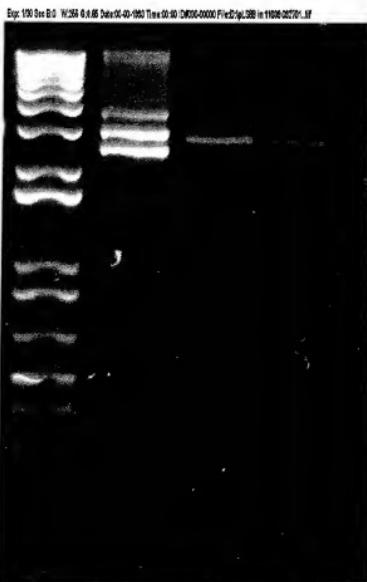
9/26/01 Wednesday - Gone to Memphis

ML started 5ml culture of ~~R6E12~~ R6E12 Ed 9 +

9/27/01 Thursday

Did a Qia spin mini prep of R6E12

Eluted in 30ml EB

1 ul
1 kb
plusR6E12
1 ulPLS88
in
P. multivittata
1029

R6E12 looks good
Pass to a new plate
freeze next week

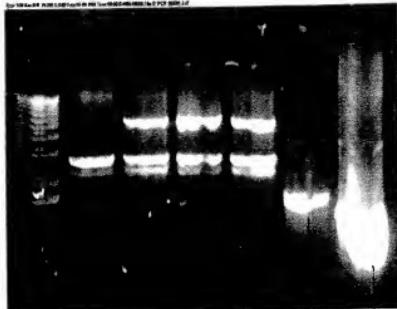
PLS88 in 1029 looks
good -

it work - we were
able to electroprorate
into *P. multivittata*
Comp cells

PCR pCLpm3 T7P1 + CLdamM20 092801

| Sample # | 1 | 2 | 3 |
|--------------------|--------|------|-----|
| pCLpm3 8/29/01 BYR | 0.5 | 1 | 0 |
| dNTPs | 5 | 5 | 5 |
| T7P1 + CLdamM20 | 1 | 1 | 1 |
| Taq | 0.5 | 0.5 | 0.1 |
| Taq Buffer | 5 | 5 | 5 |
| Water | 38 | 37.5 | 38 |
| TOTAL Rx amt. | 50 | 50 | |
| Annealing temp | 67.5 C | | |

10/2/01



see pg 41 pcr
Set up

pclpm3
lue

Load cut out of gel = 0.4177 gm
put in Ref over night

10/3/01

Elute DNA from gel with Qiagen gel extraction Kit

~~⑧~~ cut gel fragment in 1/2 and put in 2 tubes
Added 600ul / tube of QG
Added 200ul isopropanol

eluted each column w/ 30ml EB into same tube = 60ul
Called pclpm3 PCR

Loaded 5ul of pclpm3 PCR on 0.7% agarose gel 3.5ml gelstar
100 Volts



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Signed

10/3/01

Electroporated 40ul p.mutantida 11039 Comp cells w/ no DNA

Recover 1 hr in LB broth 37 C

plate 100 ul 10⁻¹, 10⁻² on LB
grow at 37 C overnight Tm = 5.16 msec
Should have plated on LB strep -
threw plates away

2.5 KV

25 sec

200 μ L

Before Digesting pCLpm3 PCR 1.7 Kb frag we need
more PCR product b/c XbaI has to be deactivated w/
EDTA (heat does not work) + then the DNA is ppted
out. DNA is lost in ppt so I need to amplify
my PCR product

| PCR pCLpm3 1.7 kb T7P1 + CLdamM20 10/3/01 | | | |
|---|--------|------|------|
| Sample # | 1 | 2 | 3 |
| pCLpm3 PCR 1.7 kb | 0.5 | 1 | 0 |
| dNTPs 1 mM each | 5 | 5 | 5 |
| T7P1 + CLdamM20 | 1 | 1 | 1 |
| Taq (7/14/01) | 0.5 | 0.5 | 0.5 |
| Taq Buffer | 5 | 5 | 5 |
| Water | 38 | 37.5 | 38.5 |
| TOTAL Rx amt | 50 | 50 | 50 |
| | | | |
| Annealing temp | 67.5 C | | |
| extension time | 30 sec | | |

run 0.5% per rxns on gel

PLS88 digest BamHI/XbaI

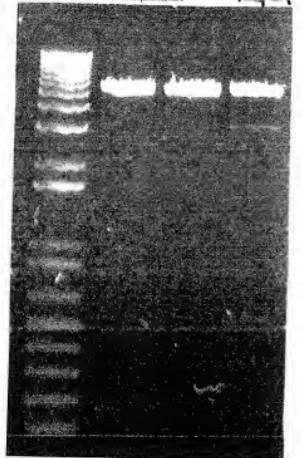
10ul PLS 8g plasmid prep 9/19/01
1ul XbaI
1ul SacI
1ul Multicore Buffer 10X
6ul H2O

200

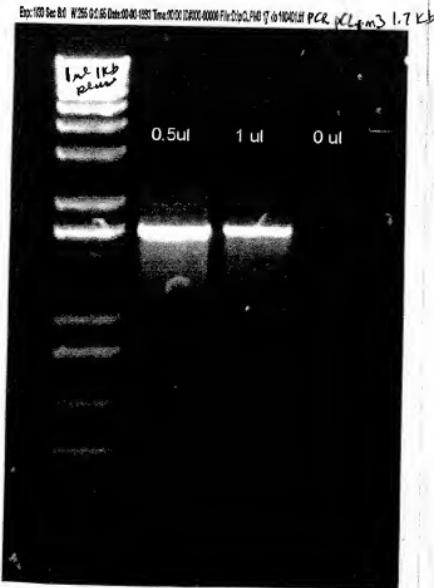
Digest 37C 4 hrs Start 10:00am

Run 0.5ml of pC1pm31.7Kb per rx run 10/3/01 on
0.7% agarose gel 3.5ml gel star

PLS38 BamHI XbaI digest
done 9/20/01



lul lul lul lul lul
1 kb BAM Kba BTRTH Xba I
plus HE



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Signed

Notebook Number:

Date:

10/4/01

Digest pCLPM3 1.7Kb frag per rxn 1 (from 10/3/01)

| | | |
|-------|----------------------|-------|
| 10 μl | DNA | |
| 1 μl | Xba I | 37 C |
| 1 μl | Bam H I | |
| 2 μl | Multicore 10X Buffer | 4 hrs |
| 6 μl | H ₂ O | |
| <hr/> | | |
| 20 μl | | |

To pLS88 & pCLPM3 1.7Kb Bam H I Xba I digests

Add 0.5M EDTA to deactivate Xba I
 Add 2 μl 3M Na Acetate (1.0 vol)
 44 μl EtOH (2 vols)

freeze -20 over night to ppt DNA.

Start 2 L BH Ap cultures of R6E12 for LPS prep

Add 2.5mls of ON culture per Liter

Add 500μl 20Dmg/ml Ap = 100ug/ml final vol
 $\frac{1}{2}$ normal conc used

grow ON 36°C Shaking

10/5/01

Cf. R6E12 cultures in 250ml bottles in G54 rotor

6000rpm for 15min at 4°C

freeze pellet at -20°C

Do LPS prep next wk

10/5/01

Cf. ppat DNA pLS88 + pCLPM3 1.7 kb
 resuspend in 10 μL TE. put at 4°C to go into
 Suspension.

10/8/01

run 1ul pLS88 + pCLPM3 1.7 kb on gel
 ✓ on gene spec first

| Sample | Type | 230nm | 260nm | 280nm | 320nm | ratio | Concentration | Dilution | Factor | Purity |
|--------------------|-------|-------|-------|-------|-------|---------|---------------|----------|--------|--------|
| pCL PM 3 1.7 kb | dsDNA | 0.655 | 1.536 | 0.818 | 0 | 1.87775 | 0.0768 ug/μL | 1 | 50 | 104.32 |
| | dsDNA | 0.707 | 1.592 | 0.85 | 0 | 1.87294 | 0.0796 ug/μL | 1 | 50 | 104.05 |
| pLS88 Bam HI Xba I | dsDNA | 1.565 | 3.982 | 2.369 | 0 | 1.68088 | 0.1991 ug/μL | 1 | 50 | 93.38 |
| | dsDNA | 1.987 | 4.493 | 2.85 | 0 | 1.57649 | 0.22465 ug/μL | 1 | 50 | 87.58 |

pCLPM3 76.8 ng/μL pLS88 211.9 ng/μL

Ligation

pLS88 4.5 Kb
 pCLPM3 1.5 Kb

pLS88 1ul
 pCLPM3 7.5ul
 Ligase Buff 1ul
 Ligase 0.5ul
10ul

14°C overnight

10/9/01 Tuesday

Heat inactivate ligation 65°C 15'

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Signed

PCR 93146 EI OPS primers 10/17/01

| Sample # | 1 | 2 | 3 | 4 |
|---------------------------------|------|------|------|------|
| 93146 genomic prep 7/28/99 | 2 | 2 | 2 | 2 |
| dNTPs 2.5mM | 10 | 10 | 10 | 10 |
| EI OPS PCR U 2 + EI OPS PCR R 2 | 1.2 | 1.2 | 1.2 | 1.2 |
| EXL polymerase | 1 | 1 | 1 | 1 |
| DMSO | 0 | 0.5 | 1 | 1.5 |
| Stabilizing Soln | 1 | 1 | 1 | 1 |
| 10 X Buffer | 5 | 5 | 5 | 5 |
| Water | 29.8 | 29.3 | 28.8 | 28.3 |
| TOTAL Rx amt. | 50 | 50 | 50 | 50 |
| Use PCR EXL Pcr Program | | | | |
| Annealing Temp | 63 | | | |
| PCR EXL primer | | | | |

Ran out of EXL polymerase after 1st rx.

ran 1st rx ordered more EXL poly

Rxs 2+ are in green RR box -20 chest freezer

gel picture pg 54

Make 2 5ml LB Ap cultures of pFPV25 w/ DH10B^S

- 1 - plasmid prep
- 1 - freeze

Make 5ml culture of pLS88 isolates pos + Adj on LBstrep positive colony from ~~better~~ adjacent to hybridization positive colony

Read and Understood By

Signed

Signed

Date:

10/18/01

Started cultures in wrong antibiotics
Started pFPV25 in LB strep. pLS88 in LB Ap opps -

Started new 5 ml cultures
pFPV25 in LB Ap
pLS88 in LB Strep

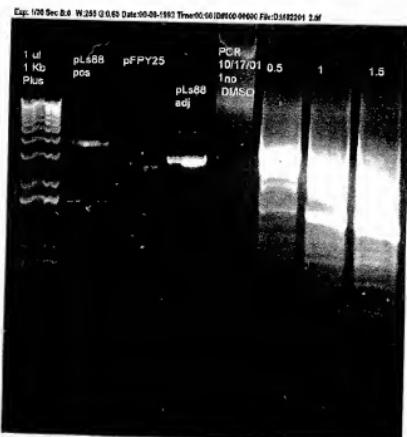
10/19/01

Froze 1 pFPV25 culture (Ask B/Y where it is)

Spin down other pFPV25 culture + both pLS88 cultures
(pos + adj) - freeze pellets for plasmid prep

10/22/01

Made QiaSpin plasmid prep of
pFPV25, pLS88 positive. pLS88 adjacent



Loaded 1ul of
pLS88 pos
pLS88 adj
pFPV25

Loaded 5ul of
pCL produced

pass both pLS88 colonies
to a new LB Strep plate

Read and Understood By

Digest pLS88 pos + Adj plasmid prep (10/22/01)

Culture
insert

| | | | | |
|------------------|-------|-------------------|-----------|------|
| pLS88 pos | 4 ul | → | pLS88 Adj | 4 ul |
| Bam HI | 1 ul | | | |
| Xba I | 1 ul | | | |
| Mnecotore | 2 ul | | | |
| H ₂ O | 12 ul | | | |
| | | 20 ul | → | |
| | | Load 10 ul on gel | → | |

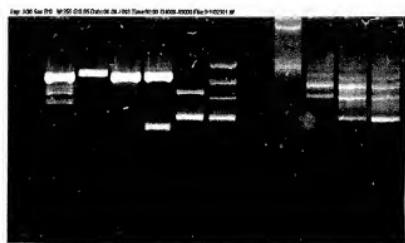
BamHI / XbaI cuts
the insert out

library

| | | | | |
|------------------|------|-------------------|------------------|-------|
| pLS88 pos | 2 ul | → | pLS88 Adj | 1 ul |
| EcoRV | 1 ul | | EcoRV | 1 ul |
| Buff D | 1 ul | | Buff D | 1 ul |
| H ₂ O | 4 ul | | H ₂ O | 1 ul |
| | | 10 ul | | 10 ul |
| | | Load 10 ul on gel | → | |

library

| | | | | |
|------------------|------|-------------------|------------------|-------|
| pFPV25' | 5 ul | → | pFPV25' | 5 ul |
| EcoRV | 1 ul | | pst | 1 ul |
| Buff D | 1 ul | | Buff H | 1 ul |
| H ₂ O | 3 ul | | H ₂ O | 3 ul |
| | | 10 ul | | 10 ul |
| | | Load 10 ul on gel | → | |



↓
1 kb plus
did not show
up

- 1) pLS88 adj BamHI/XbaI
10 ul + 2 ul 6X Dige
 - 2) pLS88 pos EcoRV
 - 3) pLS88 Adj EcoRV
 - 4) pLS88 pos BamHI/XbaI
 - 5) pFPV25' pst 1
 - 6) pFPV25' EcoRV
 - 7) PCR 10/17/01: No DNA 2 ul H₂O + 3 ul H₂O
- 8 ↓ 0.5 ul
9 ↓ 1 ul
10 ↓ 1.5 ul

Read and Understood By

Signed

Signed

Date: _____

10/24/01

| PCR pMBE1 E1 OPS primers 10/24/01 | | | |
|--|------|------|--|
| Sample # | 1 | 2 | |
| From 9/7/00 in λ E.1 pMBE1 DNA Box | 0.5 | 0.5 | |
| dNTPs 2.5mM | 10 | 10 | |
| E1 OPS PCR U 2 + E1 OPS PCR R 2 | 1.2 | 1.2 | |
| EXL polymerase | 1 | 1 | |
| DMSO | 0 | 0.5 | |
| Stabilizing Soln | 1 | 1 | |
| 10 X Buffer | 5 | 5 | |
| Water | 31.3 | 30.8 | |
| TOTAL Rx amt. | 50 | 50 | |
| Use PCR EXL Pcr Program | | | |
| Annealing Temp | 63 | | |
| PCR EXL primer | | | |

pLS88 positive gene spec

| Type | 230nm | 260nm | 280nm | 320nm | ratio | Concentration | Dilution | Factor | Purity |
|-------|-------|-------|-------|-------|---------|---------------|----------|--------|--------|
| dsDNA | 3.461 | 5.406 | 4.733 | 0 | 1.14219 | 0.2703 ug/uL | 1 | 50 | 63.46 |
| dsDNA | 0.046 | 1.798 | 1.112 | 0 | 1.61691 | 0.0899 ug/uL | 1 | 50 | 89.83 |
| dsDNA | 0.018 | 1.781 | 1.091 | 0 | 1.63245 | 0.08905 ug/uL | 1 | 50 | 90.69 |
| dsDNA | 0.031 | 1.779 | 1.083 | 0 | 1.64266 | 0.08895 ug/uL | 1 | 50 | 91.26 |
| dsDNA | 0.3 | 2.076 | 1.421 | 0 | 1.46094 | 0.1038 ug/uL | 1 | 50 | 81.16 |
| dsDNA | 0.732 | 2.531 | 1.855 | 0 | 1.36442 | 0.12655 ug/uL | 1 | 50 | 75.80 |
| dsDNA | 0.538 | 2.353 | 1.707 | 0 | 1.40187 | 0.11965 ug/uL | 1 | 50 | 77.88 |
| dsDNA | 0.37 | 2.2 | 1.522 | 0 | 1.44547 | 0.11 ug/uL | 1 | 50 | 80.30 |

pLS88 plasmid prep made 10/24/01 $\text{A}_{260} = 124.8 \text{ ng}/\mu\text{L}$

Sig pLS88 w/ CLPM1-PL 3.2 pmol + w/ T3 3.2 pmol
 $\xrightarrow{\text{p.29 bKV}}$ $\xleftarrow{\text{pg 27 bKV}}$

| | | |
|--------|------------------------------------|-------------|
| PLS88 | 4 μl | #1 CLPM1-PL |
| TRI | 8 μl | #2 T3 |
| primer | 2 μl | |
| H2O | 6 μl | |
| | <u>20 μl</u> | |

Sequence should contain Dam gene sequence.

CLPM1-PL primer sequence data is trashy - but

I blasted a segment of it and it is the Dam gene sequence

Redigest pFPV25 5 μl

| | | |
|--------------------|---------------------------------------|-------------------------|
| Buff H | 1 μl | |
| H2O | 3 μl | |
| EcoRI | 1 μl | Pst I 1 μl |
| Balgase | 1 μl | |
| | | 10 μl digest |

pFPV25 5 μl

| | | |
|--------|------------------------------------|--|
| EcoRI | 1 μl | |
| Pst I | 1 μl | |
| Buff H | 2 μl | |
| H2O | 1 μl | |
| | <u>20 μl</u> | |

Read and Understood By

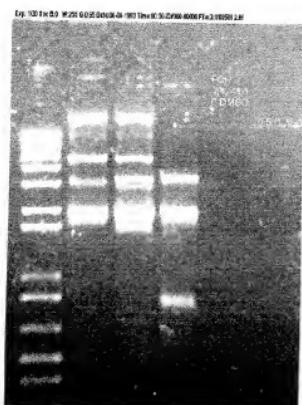
Signed _____

Signed _____

Notebook Number: _____

Date: _____

10/25/01



Freeze pLS88 pos ~~lacZ~~
Start 5ml LB Strep

Electroporate pLS88 pos into P. mult 11039 Comp cells

2.5 KV
200 Ω
2.5 uF
Tc = 5.08ms

40ul P. mult (All D4 9/21/01).
1 ul pLS88
Recover 1 hr LB Strep

plate 10⁺¹ on LB Strep
Incubate 37 °C ON

10/26/01

Exhibit A Page 46

Make QiaSpin plasmid prep of pFIV25 in λ DH10B β s

Elute 3Qul EB

Digest w/ EcoRI / Pst I (same as yesterday)

Digest 5 hrs

Run on gel w/ undigested plasmid

Froze Pls 88 ps now called pls 88 DAM

it is in XL1B MRF'

AJL D5-9 LBstrep + 20% glycerol

Put plate from Electroporation in ref over weekend -
Monday pick 4 big colonies + plate on LB strep

Started BTI Ap plate cultures of 93146 WT Lux +
93146 R4 Lux

Imaged fish immersion + ip injected w/ 93146 WT Lux
on Night Owl.



Read and Understood By

Signed _____

Signed _____

Luminous area

pLS88 positive gene spec

| Type | 230nm | 260nm | 280nm | 320nm | ratio | Concentration | Dilution | Factor | Purity |
|-------|-------|-------|-------|-------|---------|---------------|----------|--------|--------|
| dsDNA | 3.461 | 5.406 | 4.733 | 0 | 1.14219 | 0.2703 ug/uL | 1 | 50 | 63.46 |
| dsDNA | 0.046 | 1.798 | 1.112 | 0 | 1.61691 | 0.0899 ug/uL | 1 | 50 | 89.83 |
| dsDNA | 0.018 | 1.781 | 1.091 | 0 | 1.63245 | 0.08905 ug/uL | 1 | 50 | 90.69 |
| dsDNA | 0.031 | 1.779 | 1.083 | 0 | 1.64266 | 0.08895 ug/uL | 1 | 50 | 91.26 |
| dsDNA | 0.3 | 2.076 | 1.421 | 0 | 1.46094 | 0.1038 ug/uL | 1 | 50 | 81.16 |
| dsDNA | 0.732 | 2.531 | 1.855 | 0 | 1.36442 | 0.12655 ug/uL | 1 | 50 | 75.80 |
| dsDNA | 0.538 | 2.353 | 1.707 | 0 | 1.40187 | 0.11965 ug/uL | 1 | 50 | 77.88 |
| dsDNA | 0.37 | 2.2 | 1.522 | 0 | 1.44547 | 0.11 ug/uL | 1 | 50 | 80.30 |

pLS88 plasmid prep made 10/24/01 $\text{A}_{260} = 124.8 \text{ ng}/\mu\text{L}$

Seq pLS88 w/ CLPM1-PL 3.2 pmol + w/ T3 3.2 pmol
 $\xrightarrow{\text{p.29 bKV}}$ $\xleftarrow{\text{pg 27 bKV}}$

| | | |
|--------|------------------------------------|-------------|
| PLS88 | 4 μl | #1 CLPM1-PL |
| TRI | 8 μl | #2 T3 |
| primer | 2 μl | |
| H2O | 6 μl | |
| | <u>20 μl</u> | |

Sequence should contain Dam gene sequence.

CLPM1-PL primer sequence data is trashy - but

I blasted a segment of it and it is the Dam gene sequence

Redigest pFPV25 5 μl

| | | |
|--------------------|---------------------------------------|-------------------------|
| Buff H | 1 μl | |
| H2O | 3 μl | |
| EcoRI | 1 μl | Pst I 1 μl |
| Balgase | 1 μl | |
| | | 10 μl digest |

pFPV25 5 μl

| | | |
|--------|------------------------------------|--|
| EcoRI | 1 μl | |
| Pst I | 1 μl | |
| Buff H | 2 μl | |
| H2O | 1 μl | |
| | <u>20 μl</u> | |

Read and Understood By _____

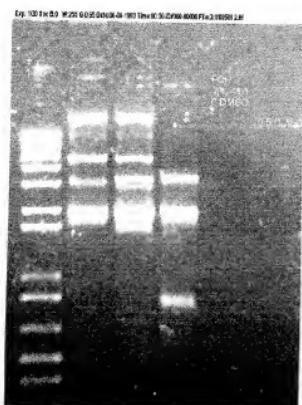
Signed _____

Signed _____

Notebook Number: _____

Date: _____

10/25/01



Freeze pLS88 pos ~~10/24/01~~
Start 5ml LB Strep

Electroporate pLS88 pos into P. mult 11039 Comp cells

2.5 KV
200 Ω
2.5 uF
Tc = 5.08ms

40ul P. mult (All D4 9/21/01).
1 ul pLS88
Recover 1 hr LB Strep

plate 10⁺¹ on LB Strep
Incubate 37°C ON

16/16/01

Exhibit A Page 49

Make Qiaospin plasmid prep of pFIV25 in DH10B_S

Elute 3Qul EB

Digest w/ EcoRI / Pst I (same as yesterday)

Digest 5 hrs

Run on gel w/ undigested plasmid

Froze pLs 88 pwo now called pLs 88 DAM

it is in XL1B MRK'

AJL D5-9 LBstrep + 20% glycerol

Put plate from Electroporation in ref over weekend -
Monday pick 4 big colonies + plate on LB strep

Started BTI Ap plate cultures of 93146 WT Lux +
93146 R6 Lux

Imaged fish immersion + ip injected w/ 93146 WT Lux
on Night Owl.



Read and Understood By

Signed _____

Signed _____

Luminous area

60 10/31/01 Pick 4 colonies from P.mut 11034 p_{BS}88
Notebook Number: _____
electroporation. grow on LB Strepate: _____

Exhibit A Page 50

11/5/01

Pour 12% / 4% SDS-PAGE gel
gel stayed in buffer in 4°C ON

Start Small LB Strep cultures of P.mut 11034 pLS 88 From 10/31
11/6/01

Samples to run on gel

LPS from R6E12 dilute - 1/1 w sample Buffer

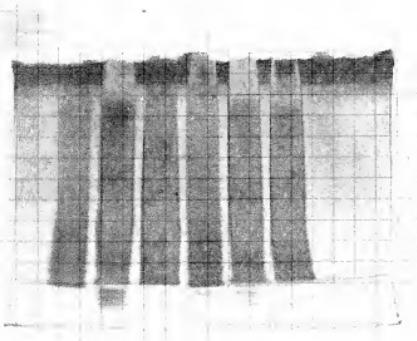
and take 10ul (1mg/ml) + 15ul H₂O + 25ul Buffer = 10.4ug

Lps Salmonella typh 1/1 of 2mg/ml in H₂O 1/1 : Buffer

Lane 1 Sal typh
2 R6E12 1/1 } Not Denatured
3 R6E12 10ug }
4 Sal typh
5 R6E12 1/1 } Denatured
6 R6E12 10ug }

loaded 10ul / well Ran at 100volts

Silver stain for LPS



Made Qia spin plasmid preps of P.mut 11039 pL588
+ colonies

11/7/01

Run 12% / 4% SDS PAGE gel

Sent ass 11/6/01

- 1) Sol. type 1/1 of Zmgluc in H2O 1/1 in Buffer - Load 10ul
- 2) R6Eiz 1/1 in buffer
- 3) R6Eiz 10ug
- 4) wt 93146 LPS 10ug

Electrophoresed at 100V/hrs

transferred to Nitrocellulose 100 volts 1 hr

Blocked Membrane overnight 5% NFM in PBS

Our Ed 9 was contaminated - so was Bobbie's

DH10Bsr pFPV25 frozen -80C All F1-5

tube w/ used to start a 5ml LBap culture

* put on LBap plate

P.mut 11039 pL588 #1 Start 5ml LB strep culture to
freeze → see off pic next page

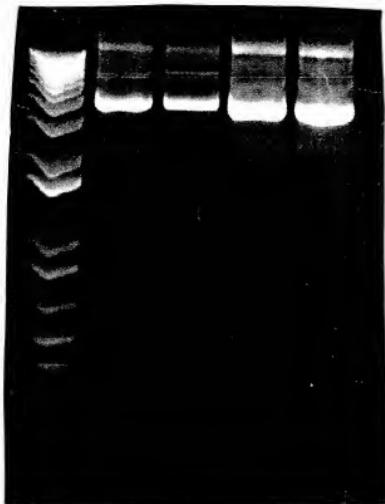
Read and Understood By

Signed

Signed

Notebook Number:

Date:

1 kb
Kopfus

11/8/01

Western Blot R6E12

There is No Ed⁺ to use as 1st Ab so

1° used α Ed^{ict} 4383 polyclonal CF serum 1/1600
+ α Ed^{ict} 93146 wt polyclonal CF serum 100μl added
to 1/100 dilution

Incubate RT 1 hr shaking
wash 3X PBS TW20

2° GEL 1/4 1 hr RT shaking
wash 3X PBS TW

p. mult 11039
Dan gene plasmid
plasmid preps
from 11/6/01

3^o goat mouse Ig Ap 1/1500 1 hr RT shaking
 Develop BC1B/NBT 20 min

it did not work
 will have to repeat
 when Bobbie has more
 Ed 9

1 2 3 4

Froze *pastuella multocida* 11039 pLS88 (DAM gene)
 LB strep + 20% glycerol
 -80°C Box A II G1-5

Digest pLS88 p.mult 11039 #1

| | |
|-----------------------------|----------------------------|
| BamH1/XbaI | |
| DNA 4 μ l | BamH1 |
| BamH1 1 μ l | DNA 4 μ l |
| XbaI 1 μ l | BamH1 1 μ l |
| Multicore 1 μ l | Multicore 1 μ l |
| H ₂ O 12 μ l | H ₂ O 4 μ l |
| <hr/> | |
| 2ml | |

| | |
|----------------------------|-------|
| | BamH1 |
| DNA 4 μ l | |
| BamH1 1 μ l | |
| Multicore 1 μ l | |
| H ₂ O 4 μ l | |

Digest 37°C
 for 2 hrs approx
 run on gel tomorrow

Plasmid prep pFV95 in DTT/DBS
 Eluted in 30 μ l set next day for gel pic

Read and Understood By

Signed

Signed

11/8/01

Notebook Number:

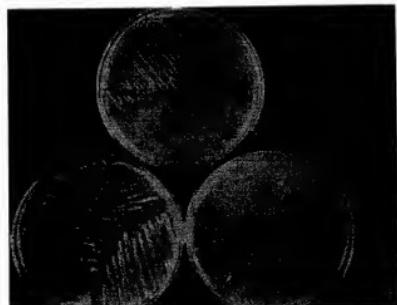
Date:



pFPr25

do not know what this band is)

Next run orange and extract top band + elutriate into $\chi L1B$



93146 Lux
recovered from
fish immersion
infected

Bacteriawell
Cultured
4 days
post
immersion

Lux~~+~~^{Plasmid}
is still
stable
in VIVD

Neg Contests
14 + 15

22: 93146 Lucy

1960-1961
2000-2001
2001-2002

Board and Standard Dev.

PCR pCLpm3 1.7 kb T7P1 + CLdamM20 purified
01/16/02

| Sample # | 1 | 2 |
|----------------------------|--------|-------------------------------|
| pCLpm3 PCR 1.7 kb purified | 0.5 | 0 |
| dNTPs 1mM each | 5 | 5 |
| T7P1 + CLdamM20 | 1 | 1 |
| Taq (7/16/01) | 0.5 | 0.5 |
| Taq Buffer | 5 | 5 |
| Water | 38 | 38.5 |
| TOTAL Rx amt. | 50 | 50 |
| Annealing temp | 67.5 C | program melt under mark |
| extension time | 30 sec | |

1/18/02

run SstI on gel of PCR Rx

SstI DNA
Inv 6x Dye

measured pMBRp4 on gene spec - got a low reading of 11.8ng/ul
So ran 1ul on gel -
Conc on gel looked like = 100ng/ul

digest w/ BamHI HindIII

| | 4 ul | 4 |
|------------|--------|----------|
| pMBRp4 | 4 ul | 4 |
| enzyme | 0.5 ul | 0.5 each |
| 10X Buff E | 1 ul | 1 |
| H2O | 4.5 ul | 4 |

digest from 2:00-5:00 37°C

Read and Understood By

Signed

Signed

2/13/02.

Ligation To End Conversion Rx mix add:

- (1) Lul plsgg vector (pls vs EcolI digest)
Lul ligase
 - (2) + controls
Lul pT7 Blue Vector
Lul ligase
 - (3) - Control
Lul pT7 Blue Vector
Lul ligase
- 22°C. 15"
- ~~22°C. 15"~~

Transformation:

Add Lul ligation Rx to 1 tube of Nova Blue Comp cells

Incubate on ice 5 min

heat shock 30 sec in 42°C water bath

Incubate on ice 2 min

Add 250ul RT SOC Media

plate

(+) + (-) Controls 50ul = 1/10 (5ul in 45ul H₂O)

on LB S-gal

exp. 50ul = 1/10 on LB strep

2/14/02

Colonies from ligation were very small first thing this AM.

Read and Understood By

Signed

Signed

Notebook Number: _____

Date: _____

3/14/02 Thursday

| PCR pCLpm3 1.7 kb T7Sal I P2 + CLdamECoRV 3/14/02 | |
|--|--------|
| Sample # | 1 |
| pCLpm3 PCR 1.7 kb purified (1/23/02) | 0.5 |
| dNTPs 1mM each | 5 |
| T7Sal IP2 +CLDamECoRV | 1 |
| Taq (7/16/01) | 0.5 |
| Taq Buffer | 5 |
| Water | 38 |
| TOTAL Rx amt. | 50 |
| | |
| | |
| Annealing temp | 55 |
| extension time | 30 sec |

used wrong template
 repeat using
 pCLpm3 plasmid prep
 8/29/01 red box G8

15 7380-085 Mar 7 2002
 GINDS15
 CLDamECoRV
 5'-CGTTTGTATGATGTTCTGTT
 11 bp
 341.8 bp
 16 KDa
 61.2 mol

15 7380-085 Mar 7 2002
 GINDS15
 T7SP1P2
 5'-GGGTGATAGCTTGATGAA
 11 bp
 584.3 bp
 15 KDa
 65.1 mol

reconstituted primers
 2.5ul each + 95ul H2O

Made new 1mM dNTPs

40ul 10mM mix + 60ul H2O

plated pLSS8 from frozen stock A3 E4 put on LB streak at 37C MLL will take out of incubator + put at 4 C tomorrow

Notebook Number: _____

Date: _____

effluvia

Hybridize colonies from Missy Person's ligation
follow protocol pg. 50

primes

PCR prod green PCR box A9 - PCR Rx 3/21/02
pg. 94 bRT = 535 ng/ul

Cleanup PCR prod using microcon columns -
follow directions on pg.

Removes primers & dNTPs

No Read on Gene Spec -

| Type | 230nm | 260nm | 280nm | 320nm | ratio | Concentration | Dilution | Factor |
|-------|-------|-------|-------|-------|---------|---------------|----------|--------|
| dsDNA | 0.709 | 0.366 | 0.224 | 0 | 1.63393 | 0.0183 ug/uL | 1 | 50 |
| dsDNA | 0.736 | 0.655 | 0.497 | 0 | 1.31791 | 0.03275 ug/uL | 1 | 50 |
| dsDNA | 0.492 | 0.945 | 0.804 | 0 | 1.17537 | 0.04725 ug/uL | 1 | 50 |
| dsDNA | 0.551 | 0.997 | 0.861 | 0 | 1.15796 | 0.04985 ug/uL | 1 | 50 |

There are crazy readings - gene spec is
not good for measuring DNA of PCR prod.

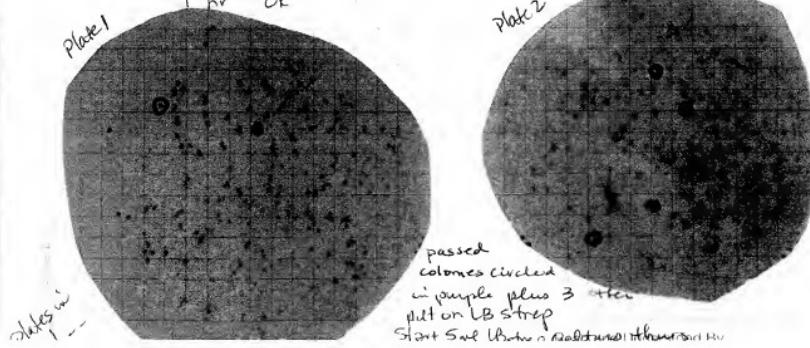
(MLL and ML3 bulk w/ probe)

Remained of cleaned up PCR prod is in pink box 118

Hybridize ON 42°C

4/17/02 Developed Hybridization

- 1" film almost had no images
- 10" very light image
- 1 hr OK



5/13/02

cycle seqy PGEM 3Z test plasmid w/ M13 primer from Ab1
run on Seq. to create matrix file

5/14/02 Tues

Clones from Electroporation grew well

grow 2ml LB strap cultures
pick colonies w/ sterile toothpick
use 1ml tomorrow to do plasmid preps for insert

grow 1 culture from Neg Control

| | | | |
|-----------------|--------|-------|----|
| pl588 pCPM3 | 100 μl | = 150 | 53 |
| | 50 μl | 26 | 5 |
| Neg Con (pl588) | 100 μl | 21 | 55 |
| No insert | 50 μl | 14 | 11 |

pick 10 ligation colonies
1 neg con colony



5/15/02 Wed

All 2 ml cultures grew

used 1ml to do Qiaprep spin plasmid preps

eluted in 30ml EB

Run 3ml DNA on gel
1 μl GEX Dye
2 μl H2O
6 μl

It Finally
Worked !!

#5 & Neg
plasmid preps
in yellow box
A5 + A6

For Vaccine trial:

plated 93146 R6 from 6/19/01 A4G5 -80C on Blood
grow at 26°C 48hrs

Reconstituted Interpet Edict vaccine - filled bottle 2/3 full of sterile H2O

plate 1 loop full on Blood grow at 26°C 48hr
put 50ml into 5ml BH1 to freeze back grow at 26°C shaking
over night

1 TSDmE flask BH1 for Interpet Vaccine
4 TSDmE flask BH1 for R6

Read and Understood By

Signed

Michelle Sc

Notebook Number: _____

Date: _____

For Vaccine trial cont:

Took 5mls of Heat Killed Ed ret (BVR's prep 4/12/02)

Tared tube empty

add 5mls

Cf 12000 rpm 10"

Aspirate Sup

Re weigh tube = 0.0499 gm

Lyophilize overnight re weight

resuspended pellet in 2mls H₂O

transfer to a pre-washed

15 ml tube = 6.7018 gm

Freeze

Lyophilize

15 ml tube Dry wt = 6.6064

Dry wt of pellet = .0046

Digest plasmid + clone (#5) + Neg plasmid preps

w/ Sal I + Cla I single digests

2 μl DNA

1 μl Buffer

0.5 μl Enzyme

6.5 μl H₂O

10μl digest

ABI ~~matrix~~ matrix stds came in. stds are 4 colors: Blue, Red, Green, Yellowmixed 1μl of each Standard with 12μl of deionized formamide
(borrowed from Hanson's Lab). 1 tube per Std = 4 tubes / each color

Heat to 95°C for 2 "

Chill on ice

Sequence using Seq Run (250 μl) E run module

injected each sample 3 times - this is done when the injection list is filled out

5/10/02 Thurs

Sbs are still running on seq.

Weighed lyophilized pellet of heat previous page

Run digest on gel plasmid prep clone #5

Load entire Digest

6K dye is really conc. So only use 0.7% agarose gel

10ul DNA digested

1ul 6K Dye

1ul H2O

into 11039 comp cells All D1-3
BHI strep plate 2 on 2 100ul

Ecoli 93146 #19 pass from MW plate

0.7% agarose gel 2 more 750 ml broth cultures

run 100 Volts

load on gel

| | |
|--------|-----------------------|
| Lane 1 | lul 1Kb |
| 2 | Neg C _{la} I |
| 3 | Neg S _{aI} I |
| 4 | #5 C _{la} I |
| 5 | #5 S _{aI} I |



Electroporate: pls dam2 into 11039 p.mut. electrocompetent cells

1ul plasmid prep #5 (yellow box) A8
40ul 11039 Comp cells box All D2

Tc = 5.12

5.5 KV

200 S2

25 uF

Recovered 1 hr in LB 37C

plate 100ul neat on 2 LB Strep plates

Made 2 more 750 ml broth cultures to grow 93146 wt #19

to heat kill for vaccine trial

Read and Understood By

Lubell S
Signed

$$\frac{37g}{\text{Strept}} \times \frac{750\text{ml}}{\text{Strept}} = 27.75 \text{ gms BHI}$$

5/16/02 Thurs.

Froze

pls Dam 2 5vials 1ml/each in LBgprep + 20% glycerol
 Aquavac-ESC 5 vials 1ml/each in BH1 to 20% glycerol
 Vaccine

5/17/02 Fri:

Staff Appreciation Day:

Hofhory has grown yet on plates from Electroporation -

maybe need to be on BH1 Strep
 Maybe 11039 Comp cells Dead

Leave at 37 C until Tomorrow

Passed 93146 WT #19 to new blood plate

Start 5 ml culture Sun

Took Rec + vaccine plates out of incubator
 Leave at RT

Start 5 ml BH1 Cultures Sat.

5/19/02 - Started 5 ml BH1 Cultures

5/20/02 MM leave Mark Sick BYA Started 2 750ml cultures
 by ~~WT~~ WT #19

5/21/02 93146 WT Ed.ict take 1ml from each 750ml culture
 Combine in 1 tube

Streak for purity - plate is pure

plate 10^{-5} + 10^{-6} for colony counts

10^{-5} TNTC TNTC

10^{-4} 286 287 Avg 286.5

Heat Kill at 60°C - for 3 hours
 plate for viability - no growth at 48 hrs

7/24/02

Blotted plates on nitrocellulose membrane.

Block in 0.01M PBS + 5% NFM 2 hrs RT

1^o Ab used straight Ed 9 1 hr RT

3X wash PBS 0.01M + Tw20

2^o Ab goat or mouse Ig (H+L) Ap 1/1500 4°C overnightSet up Serum Killing Assay see pg 17 for plate setup
serum HI at 56°C for 30min

Read plate every 5min for 1 hr

plate 15ul of 10⁻⁴ + 10⁻⁴ on BH1 Ap

Colony Count Results (7/25/02)

| <i>E. coli</i> | 10 ⁻⁴ | 10 ⁻⁷ | assay worked per test |
|----------------|--------------------------------|-----------------------------|--|
| 27, 30, 47 | 2, 2, 3 | | 0.31 X 10 ⁹ |
| 7/26/02 WT | 10 ⁻⁴ 35, 21, 26 | 10 ⁻⁷ 2, 5, 9 | 1.82 X 10 ⁹ 3.56 X 10 ⁹ |
| TC | 10 ⁻⁴ 44, 30, 35 | 10 ⁻⁷ 0, 3, 6 | 2.42 X 10 ⁹ 2 X 10 ⁹ |

Injected mice

100ul ip. of pimelocida uOT 50, 100, 1000 2 mice per dose
 " Dam 50, 100, 1000
 " Dam 2 50, 100, 1000
 Control PBS 3 mice

Read and Understood By

Elizabeth S.

Signed

Signed

- 24 CO₂ indicates mouse was euthanized to reduce suffering

Notebook Number: _____

Date: _____

7/25/02 Mortality from mice Exp.

| | | | | | |
|--------|-----------|--------|-------|-------------------|-------------------------------------|
| 9:00am | WT 100 | 2 mice | Treat | 2 CO ₂ | Lethargic, Shivering, pilo erection |
| | WT 50 | 1 | DOA | | " |
| | Dam 100 | 1 | DOA | | " |
| | Dam 100 | 1 | DOA | | " |
| | Dam 2 100 | 1 | DOA | | " |

Notes:

2nd mouse in WT 50, it is very lethargic, doesn't respond to being handled
3rd mouse in Dam 100 is lethargic, Shivering (due to fever?)

| | | | | |
|---------|---------|-----------------|------------------------------|------------------------------|
| 11:30am | WT 100 | CO ₂ | Lethargic, Fever (shivering) | does not respond to handling |
| | Dam 100 | 1 | DOA | |
| | Dam 50 | 1 | DOA | |

Notes:

2nd mouse in Dam 50 is sick - shivering, non-respnsive, pilo erection

1:30 No morts

| | | | |
|------|--------|---|-----|
| 4:45 | WT 50 | 1 | DOA |
| | Dam 50 | 1 | DOA |
| | WT 100 | 1 | DOA |

All mice from TET Groups: WT 100, WT 50, WT 50, Dam 100 & Dam 50 are Dead

Dead mice were opened ventrally, the chest was spread open
they were stored in Sample cups w/ 10% buffered formalin

7/26/02

9:00am Dam 100: 1 DOA

All mice from TET Group Dam 100 are Dead

7/27 + 7/28 Dan Scruggs ✓ mice

Mon 7/29/02

Start WT, R6 + E. coli Luria cultures on BH4 Ag + S and BHI Broth

Mice - All TET groups are active + Alert
pn " " " "

8/28/02

Colony counts - Serum Killing Assay E. coli Lux

$$\begin{array}{r} 10^{-7} \\ \quad 5 \\ \quad 5 \\ \quad 5 \end{array}$$

$$\cancel{1.6 \times 10^{10} \text{ CFU/ml}}$$

$$\begin{array}{r} 10^{-4} \\ \quad 35 \\ \quad 51 \\ \quad 25 \end{array}$$

$$\cancel{3.7 \times 10^{10} \text{ CFU/ml}}$$

8/29/02

R₆ lux + WT Lux Colony Count plates were
Contaminated too badly to count

9/1/02 Inject mice

BYR prepared p multocida 11039 WT + Dan2 for
injections 1 xtra mouse uninjectected

5 mice / dose 9 treatments

20 - 4 doses WT 5, 10, 50, 100

20 - 4 doses Dan2 10, 50, 100, 500

5 control mice - PBS

Inject mice w/ 100ul each i.p.

use 1cc syringe with 27 1/2 G needle

mice were not set up properly - we (Dan Scruggs + I) had to

redistribute mice to 5/cage before injection - mice were stressed!
Injected mice 11:30am ✓ 1:30pm all OK ✓ 4:15pm all OK

Read and Understood By

Signed

Signed

9/6/02

✓ mice 8:00 Am

WT 5 - 3 mortalities

1 lethargic, shivering, unresponsive.

1 almost dead completely unresponsive, unable to walk euthanize w/CO₂
→ dead before I could euthanize

WT 10 - 4 morts

1 sluggish but responsive

WT 50 3 mort

2 sluggish but responsive

WT 100 3 mort

2 sluggish but responsive have pilo erection

Dam 2 500 All OK

100 all OK

50 all OK

10 all OK

PBS (-) control All OK

✓ mice 10:45 Am

WT 100 both mice are shivering, lethargic, very rapid breathing

✓ mice 2:45 pm

WT 100 breathing very rapidly unresponsive

WT 50 " " "

WT 10 very lethargic

WT 5 sick but alert

Dam 2 - all alert but not as alert as controls

✓ mice 6:20 pm

WT No change

Dam 2 No change

PBS No change

10-5-02

Actinomycetillus - plasmid's electroporation

- 1- Removed 40 μ l of actinomycetillus competent cells from nitrogen tube and added to a clean tube.
- 2- Added 1 μ l of plasmid DNA to the ~~same~~ tube.
- 3- Placed suspension in electroporation cuvette.
- 4- Set electroporator settings.
- 5- Electroporated and got a λ culture control = 5.26.
- 6- Placed cells suspended in 1 ml BHI-Na⁺.
- 7- Incubated for 1 hour at 37°C.
- 8- Diluted culture to 10⁻³.
- 9- Plated 100 μ l of 10⁰, 10⁻¹, 10⁻² on BHI-Na⁺ streak plates.
- 10- Incubated plates at 37°C overnight.

Results

10-5-02

Plate Count

| | |
|------------------|----|
| 10 ⁰ | 33 |
| 10 ⁻¹ | 8 |
| 10 ⁻² | 0 |

Notebook Number: _____

Date: _____

10-29-02Mutation Rates

- 1- Diluted XL1 Blue, XL1 dam⁺, Actin, and Actin dam⁺ to 10^{-6} .
- 2- Plated in triplicate with 10⁵ cfu of 10^5 and 10^4 .
- 3- Plated 100⁵ cfu of overnight culture in triplicate on BHI, i.f. 100⁵ cfu in BHI(NAS), & 100⁵ cfu.
- 4- Incubated at 35°C overnight.

Results

Plate Counts

XL1 B
XL1 dam⁺
Actin
Actin dam⁺

| | $\frac{10^5}{TNTC}$ | $\frac{10^4}{TNTC}$ |
|------------------------|---------------------|---------------------|
| XL1 B | 1 | 1 |
| XL1 dam ⁺ | 1 | 1 |
| Actin | 1 | 1 |
| Actin dam ⁺ | 1 | 1 |

BHI, i.f.

| | Bacterial lawn | Bacterial lawn |
|------------------------|----------------|----------------|
| XL1 | No growth | No growth |
| XL1 dam ⁺ | No growth | No growth |
| Actin | No growth | No growth |
| Actin dam ⁺ | No growth | No growth |

Note: Will have to do antibiotic titer with Actinomycete and *Escherichia coli*

10-29-02Fish Trial (24 hour)

- 1- Pooled 3 fish from each tank.
- 2- Framed eye and kidney.
- 3- Weighed, macerated, diluted to 10^{-2} .
- 4- Plated 15¹ cfu of 10^0 , 10^1 , 10^2 on EIM plates.
- 5- Incubated at 35°C for 48 hours.

- 1) Master cultures of HB101 30-96 and electroporation for plasmid prep.
 2) Master cultures of 9344a wt, rec, and EC 11229 for MLE micro.

Boiling with conjugation reagents

1. Pellet bacteria in microtiter plate.
2. Resuspended in 100µl HBSB 0.9% with Ampicillin.
3. Transferred 5µl from resuspended plate and added to 35µl HBSBamp.
4. Transferred 5µl from above dilution plate and added to 35µl HBSBamp in middle bottom assay plate.
5. Added one additional 40µl HBSB to control plate and 40µl normal serum to assay plate.
6. Incubated at RT for 1 hour.
7. Took reading on Dmax at Times 0, 30, and 60.
8. Analyzed Sabouraud Ecol.
9. Highlighted reagents were going back (50°C).

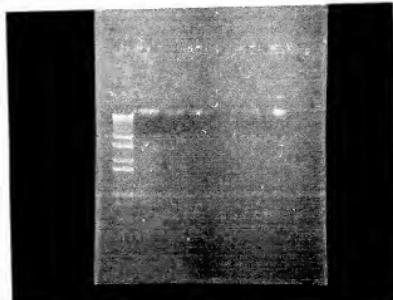
Results: See next page. (86)

1-30-93 HB101 30-96 donor plasmid prep.

1. Removed plasmid DNA with Qiagen kit.
2. Loaded 3µl on 7% agarose gel.
3. Run at 100V for 1. hours.

Lane 1 - 1 No Laddar

Lane 2 - prep 1
 Lane 3 - prep 2
 Lane 4 - prep 3
 Lane 5 - prep 4
 Lane 6 - prep 5
 Lane 7 - prep 6



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